AMENDMENT

In the Claims:

Applicant amends claim 84 and adds new claim 101 as follows:

84. (Amended) A composition [able to amplify] useful in the detection of Mycobacterium tuberculosis nucleic acid, comprising:

a) one or more oligonucleotide from about 10 to about 100 nucleotide bases in length which will, under nucleic acid amplification conditions, bind to or extend through a region of Mycobacterium tuberculosis nucleic acid consisting of a nucleotide base sequence, said region selected from the group consisting of:

[a]<u>i)</u> SEQ ID NO: 22,

[b] ii SEQ ID NO: 3,

[c] <u>iii)</u> SEQ NO: 2,

[d] \underline{iv} SEQ ID\NO: 4,

[e] \underline{v} SEQ ID NQ: 5, and

[f]vi) the nucleotide sequences perfectly

complementary to these sequences, and

b) a nucleic acid hybridization assay probe.

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- 101. (New) A composition useful in the detection of Mycobacterium tuberculosis comprising at least one oligonucleotide, or composition containing an oligonucleotide, selected from the group consisting of:
- a) a nucleic acid hybridization assay probe comprising an oligonucleotide which will hybridize to at least 10 contiguous bases of a nucleotide base sequence region of a target

 Mycobacterium tuberculosis nucleic acid, said region selected from the group consisting SEQ ID NO: 3, SEO ID NO: 8, and the sequences perfectly complementary thereto;
- b) an oligonucleotide able to bind to or extend through a region of Mycobacterium tuberculosis nucleic acid, said region consisting of a nucleotide base sequence selected from the group consisting of SEQ ID NO: 1, SEQ ID NO: 2, SEQ ID NO: 3, SEQ ID NO: 4, SEQ ID NO: 5, SEQ ID NO: 6, SEQ ID NO: 7, SEQ ID NO: 8, SEQ ID NO: 9, SEQ ID NO: 10, SEQ ID NO: 22, SEQ ID NO: 23, and the sequences perfectly complementary to these sequences,
 - c) a kit comprising the oligonucleotide of step (b), and
- d) a specifically detectable nucleic acid hybrid formed under nucleic acid hybridization conditions between the

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